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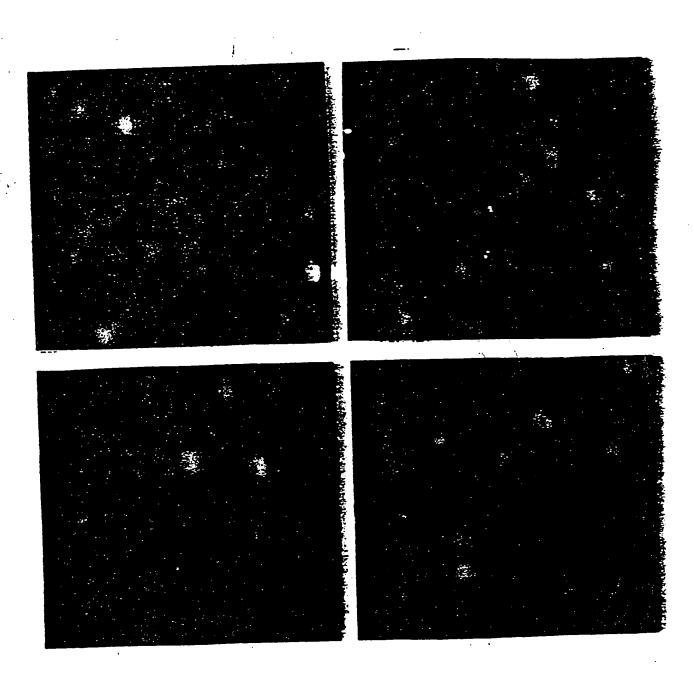


Figure 2

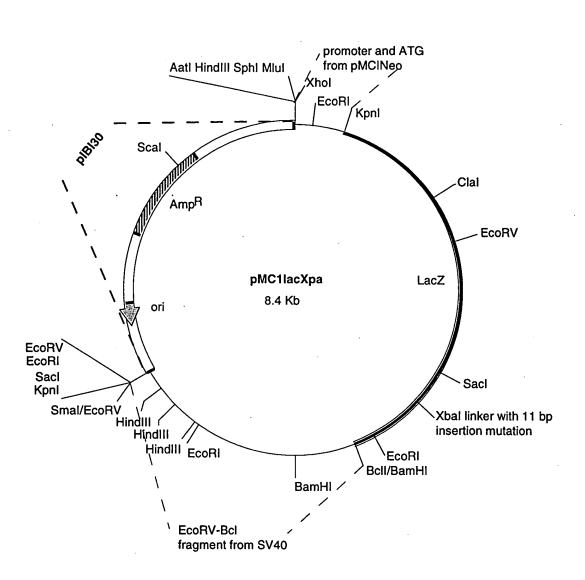


FIGURE 3

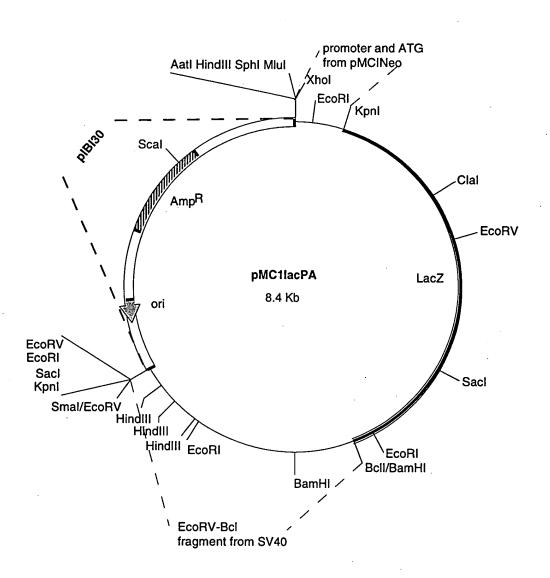


FIGURE 4

. 1

Figure 5

3610 3620 3630 3640 3650 3660 ATAAAAAACAACTGCTGACGCCGCTGCGCGATCAGTTCACCCGTGCACCGCTGGATAACG

3670 3680 3690 3700 3710 3720
ACATTGGCGTAAGTGAAGCGACCCGCATTGACCCTAACGCCTGGGTCGAACGCTGGAAGG

PCR α

3730 3740 3750 3760 3770 3780 CGGCGGCCATTACCAGGCCGAAGCAGCGTTGTTGCAGTGCACGGCAGATACACTTGCTG

3850 3860 3870 3880 3890 3900 GCCGGAAAACCTACCGGATTGATGGTAGTGGTCAAATGGCGATTACCGTTGATGTTGAAG

PCR β

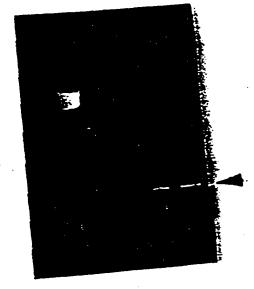
3910 3920 3930 3940 3950 3960
TGGCGAGCGATCCGGCGCGCGCGCTGAACTGCCAGCTGGCGCAGGTAG

3970 3980 3990 4000 4010 4020 CAGAGCGGGTAAACTGGCTCGGATTAGGGCCGCAAGAAAACTATCCCGACCGCCTTACTG

TEST FOR ALTERATION OF AN INSERTION MUTATION IN THE lacZ (β -GALACTOSIDASE) GENE OF EUKARYOTIC EXPRESSION VECTOR

Experimental Sample	276-mer D	d Plasmid, NA and RecA rotein	Number of Injected Surviving Cells	Number of Surviving Cells Scoring Blue	Surviving Cells Scoring Blue (%)
1 1	pSV-β-gal	-276-mer - RecA	168	21	12.5
2	pMC1lacpa	- 276-mer - RecA	98	9	9.2
3	pMC1lacXpa	- 276-mer - RecA	173	0	0
4	pMC1lacXpa	+ 276-mer - RecA	103	0	0
5	pMC1lacXpa	+ 276-mer + RecA	168	. 6	3.6

Figure 6

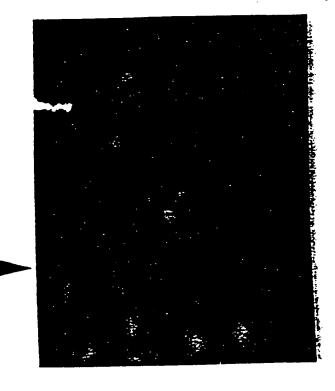




687 bp

300

1 2 3 4 5 6

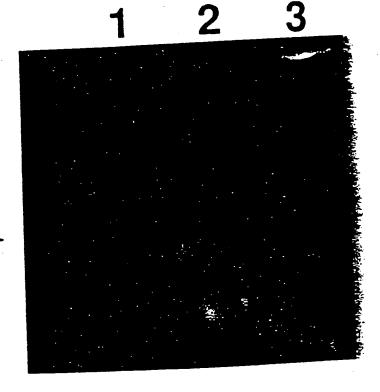


Fryre

1 2 3 4 5



FIGURE 8B



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Scheme for recombination assay

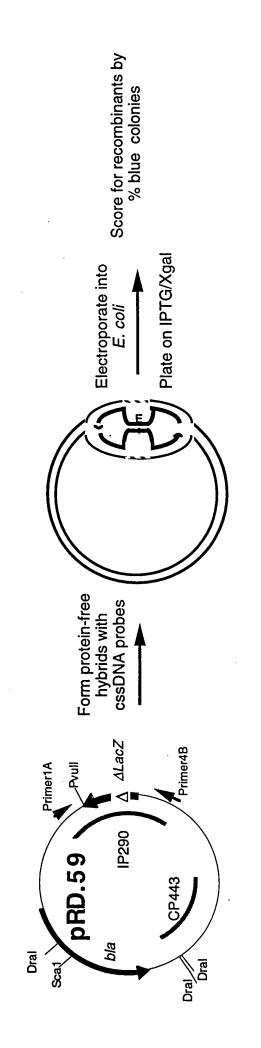


Figure 10

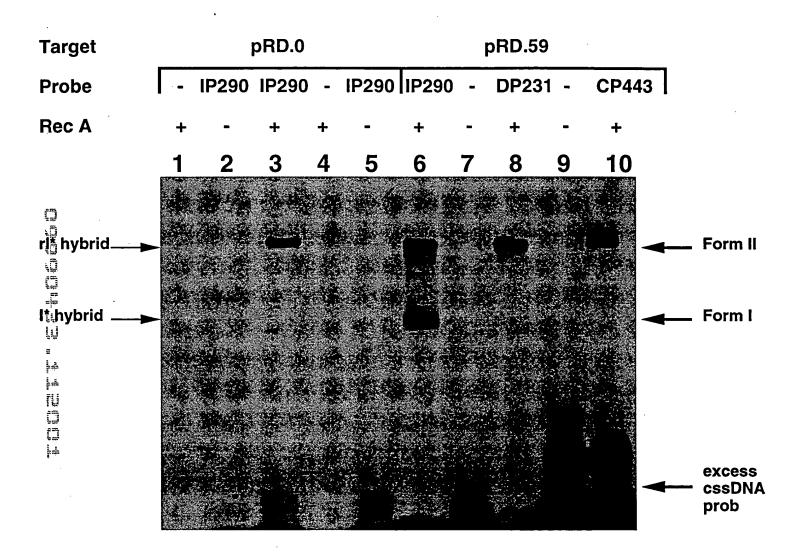


FIGURE 11

cssDNA Probe: Target Hybrids Enhance Homologous Recombination

Target Probe	RecA coating	Host	%Recombinant / total colonies
pRD.59 -	+ +	RecA+ RecA -	00
pRD.59 IP290 IP290 pRD.59 IP290 IP290	+ +	RecA+ RecA - RecA+ RecA+	0000
pRD.59 DP290 DP290 pRD.59 DP290 DP290	+ +	RecA+ RecA - RecA+ RecA+	0000
pRD.59 CP443 CP443 pRD.59 CP443 CP443	+ +	RecA+ RecA - RecA+ RecA -	0000

Figure 12

Unstable

Three-strand hybrids

Certain regions of DNA probe or target strands are unpaired

Four-strand hybrids

All DNA probe and target strands are paired

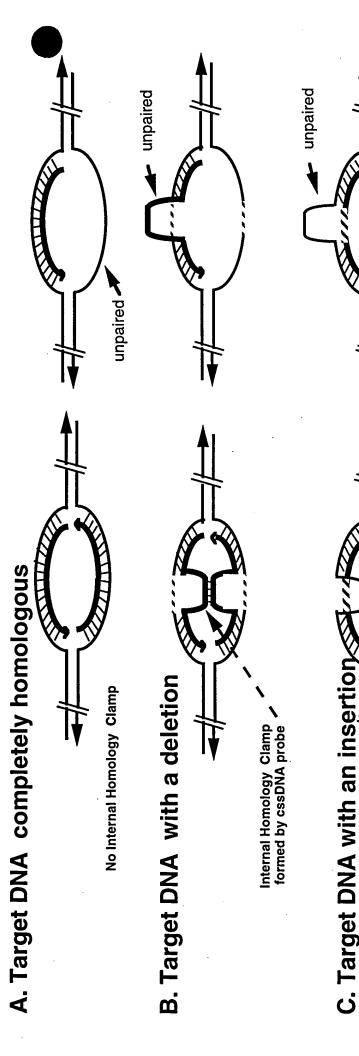
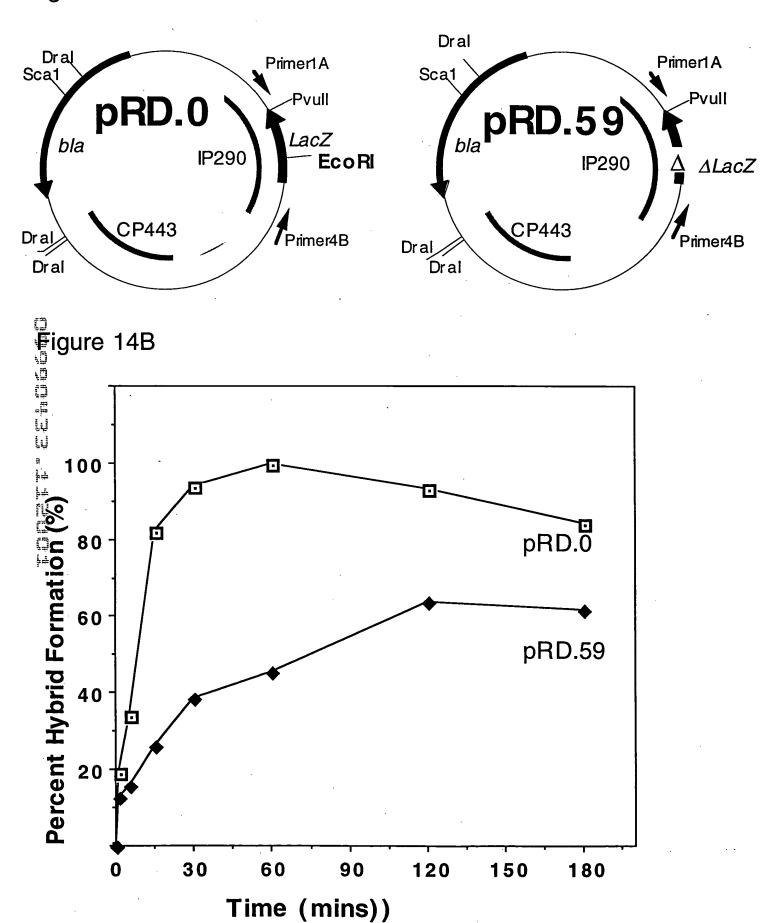


Figure 13

Internal Homology Clamp formed by dsDNA target

Figure 14A



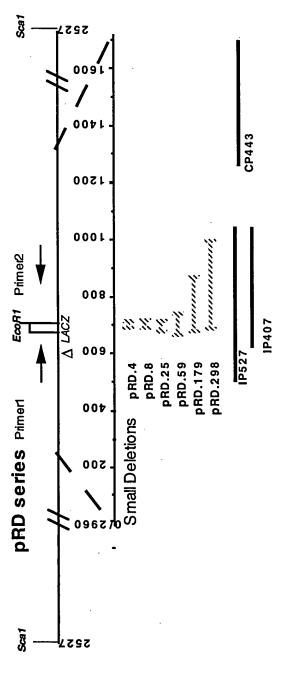


Figure 15A

 \triangleleft

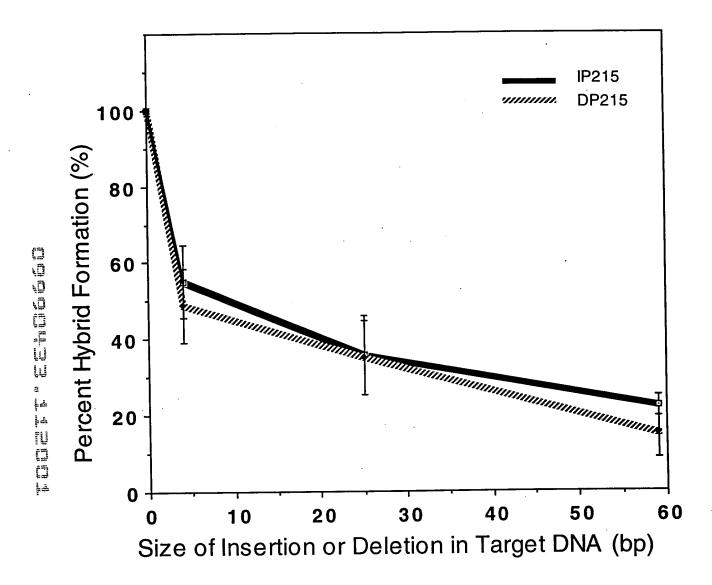
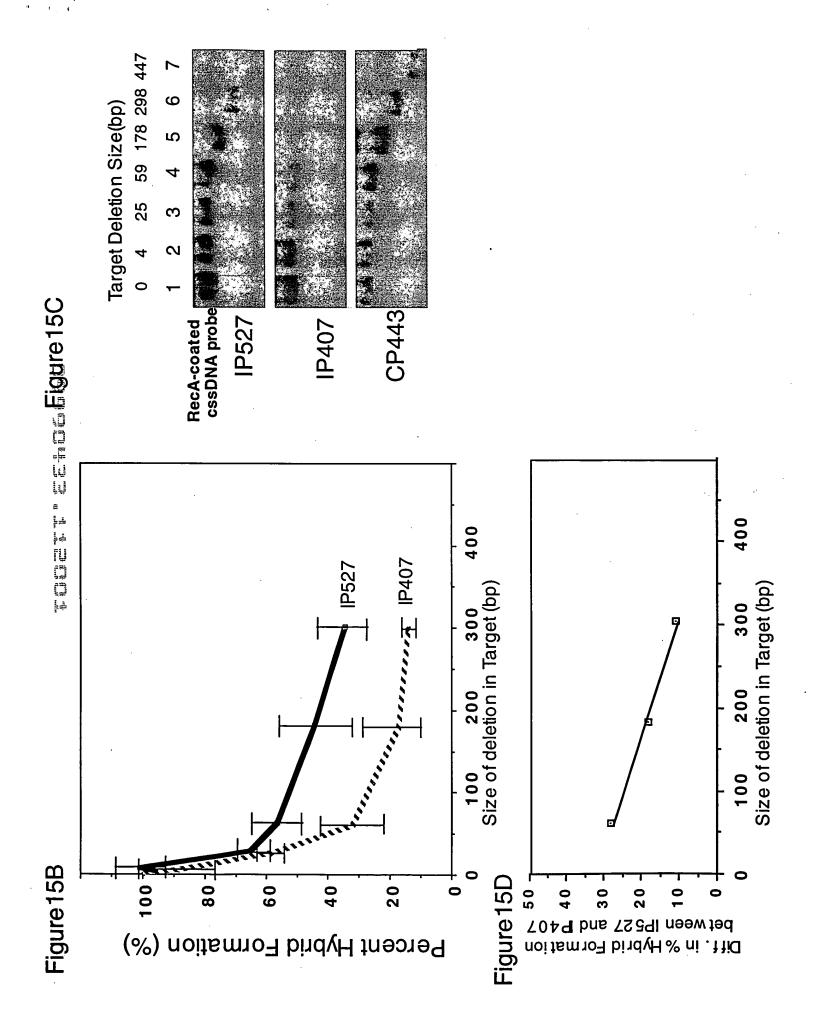


Figure 16



Function Figure 17A

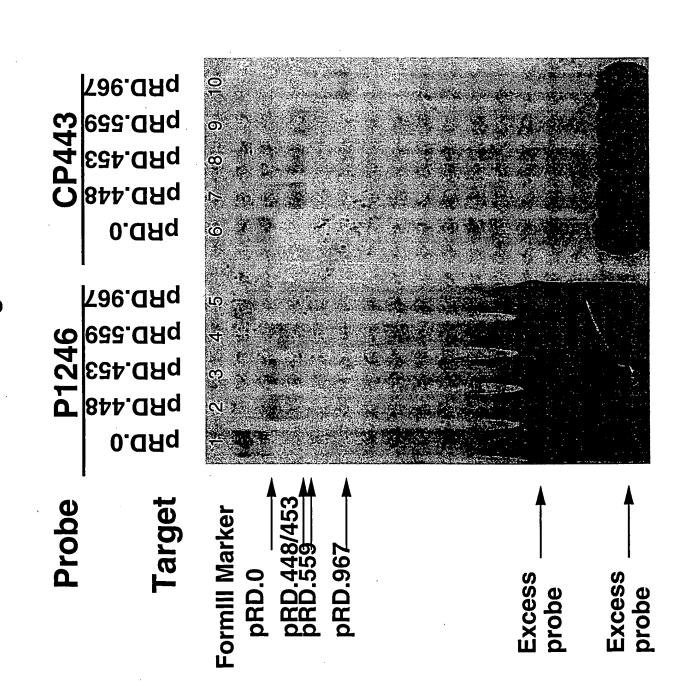


Figure 17B

HYBRIDIZATION RELATIVE HOMOL. (bp) AMT TARGET DNA

0.8 798 793 687 pRD.448 pRD.0 pRI pRI

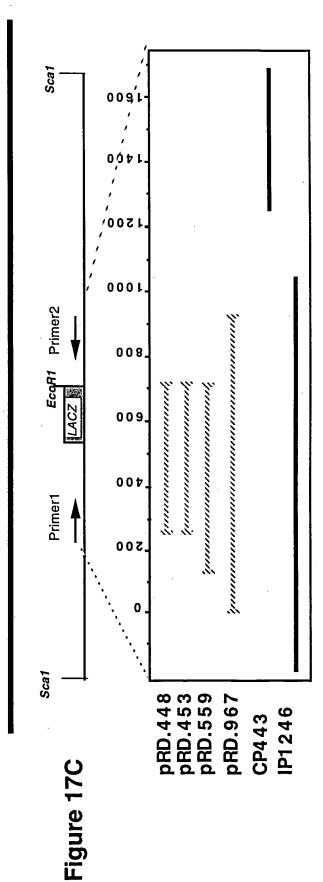
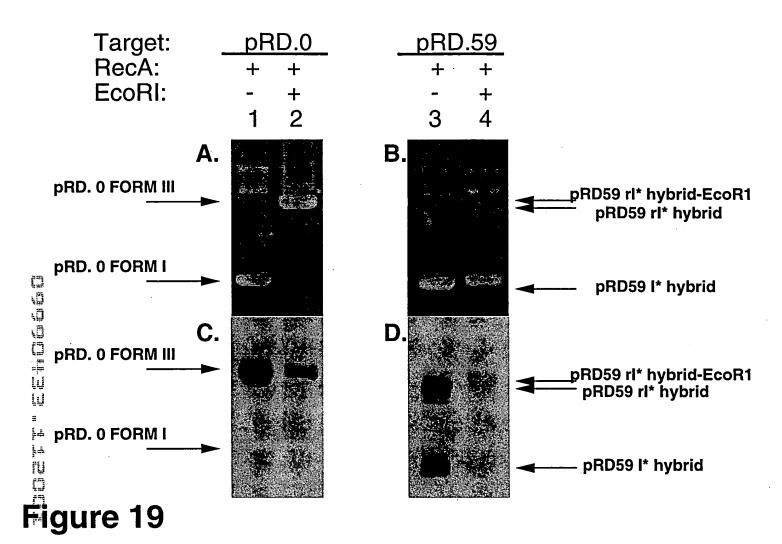
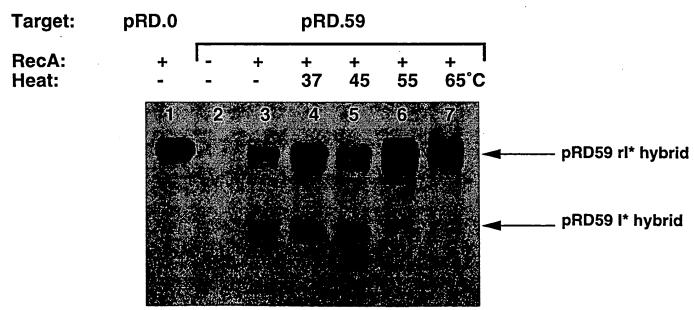
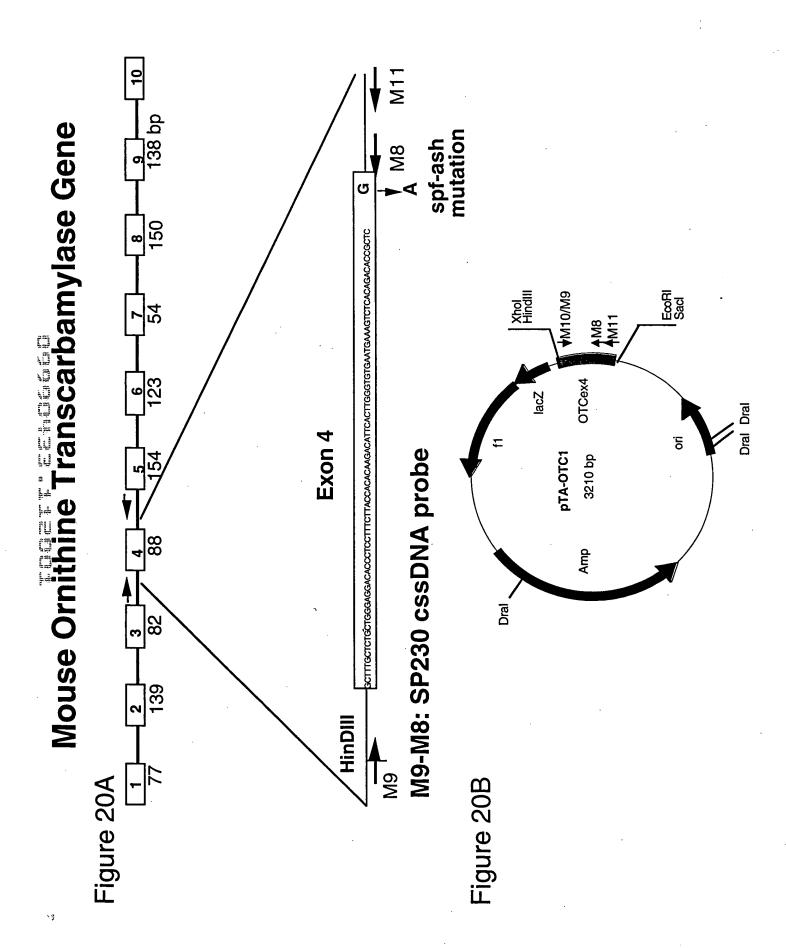


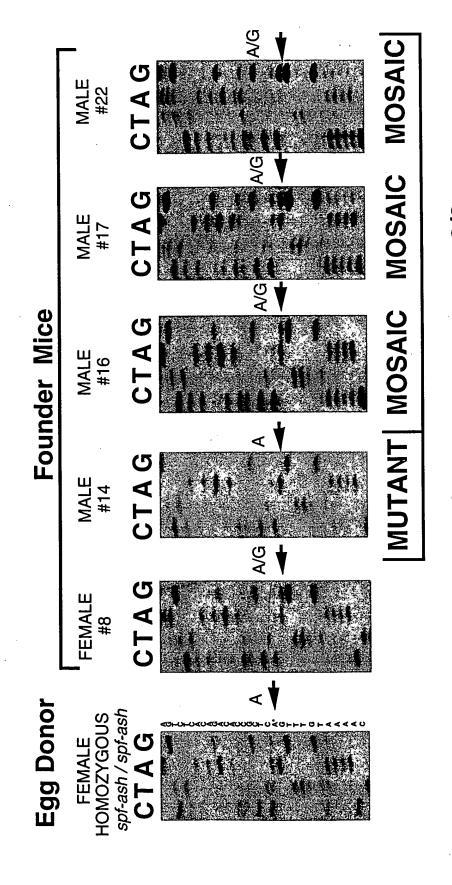
Figure 18







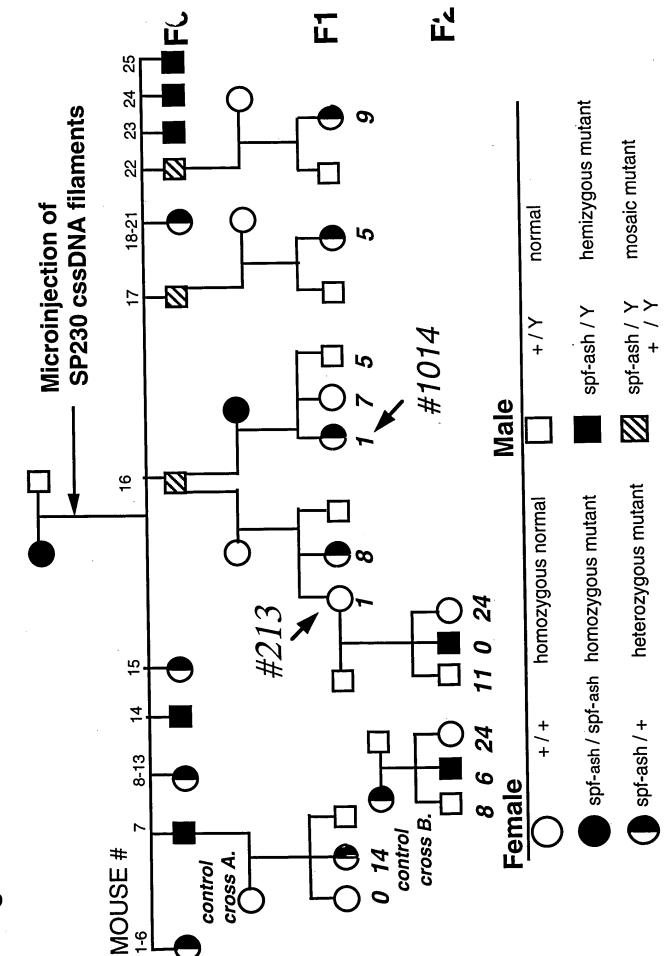
OTC exon4 DNA sequence determination

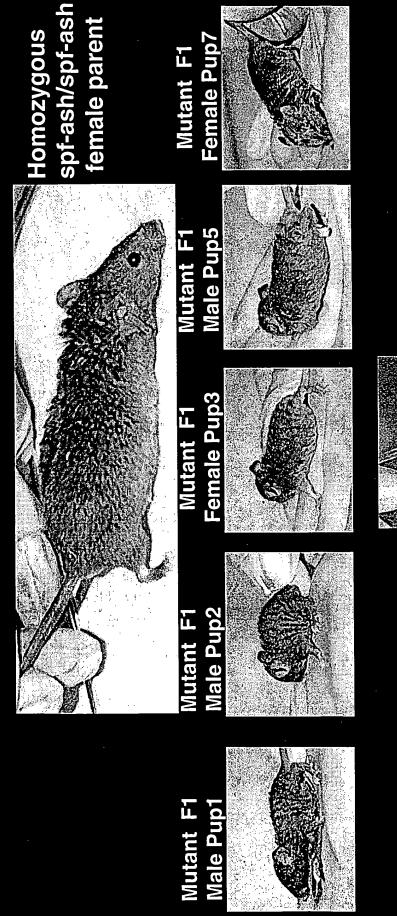


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Germline Transmission of EHR Corrected OTC+ Allele Figure 22







Normal *

F1 Female

Pup1014